

# **The Effect of $\text{Ag}^+$ on the Excited State Properties of Gas Phase (Cytosine) $_2\text{Ag}^+$ Complex: Electronic Transition and Estimated Lifetime**

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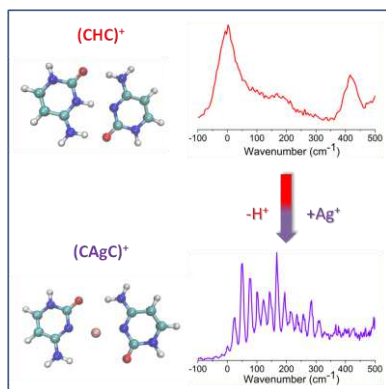
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## Abstract

Recently, DNA molecules have received great attention because of their potential applications in material science. One interesting example is the production of highly fluorescent and tunable DNA- $\text{Ag}_n$  clusters with cytosine (C) rich DNA strands. Here, we report the UV photofragmentation spectra of gas phase Cytosine... $\text{Ag}^+$ ...Cytosine ( $\text{C}_2\text{Ag}^+$ ) and Cytosine... $\text{H}^+$ ...Cytosine ( $\text{C}_2\text{H}^+$ ) complexes together with theoretical calculations. In both cases the excitation energy does not differ significantly from that of isolated cytosine or protonated-cytosine indicating that the excitation takes place on the DNA base. However, the excited state lifetime of the  $\text{C}_2\text{H}^+$  ( $\tau = 85$  fs), estimated from the bandwidth of the spectrum, is at least two orders of magnitude shorter than that of the  $\text{C}_2\text{Ag}^+$  ( $\tau > 5300$  fs).

The increased excited state lifetime upon silver complexation is quite unexpected and it clearly opens the question about what factors are controlling the non-radiative decay in pyrimidine DNA bases? This is an important result for the expanding field of metal-mediated base pairing, and may also be important to the photophysical properties of DNA-templated, fluorescent silver clusters.



**Keywords:** DNA- $\text{Ag}_n$  cluster, excited state, photofragmentation spectroscopy, DNA bases, metal-mediated base pairs

In the past years, natural and artificial DNA molecules have been of great interest because of their potential applications in biological and material science.<sup>1-5</sup> The central idea is to exchange the natural canonical interaction between the DNA bases for new non-covalent interactions (i.e hydrogen bonding, hydrophobic interactions, structure complementarity and site-specific functionalization),<sup>6</sup> leading to artificial base pairing. One of the recently established methods for site-specific functionalization is metal-mediated base pairing, that is to replace the hydrogen atom within hydrogen-bonded base pairs by metal ions.<sup>6</sup>

Metal mediated base pairing constitutes a major advance in the attempt of expanding the genetic code, at least twenty three artificial metal mediated base pairs have been reported to stabilize the double helix structure of DNA (for a complete review the readers are referred to a recent work on metal-mediated DNA base pairs that address all the studies on this topics all over the time).<sup>6</sup> Besides, the incorporation of metal mediated base pairing has proven to be a suitable and powerful tool for the potential development of artificial DNA based devices.<sup>7,8</sup>

It is recognized that the T-T and C-C mismatch pairs can be transformed into very stable T-Hg<sup>2+</sup>-T<sup>9</sup> and C-Ag<sup>+</sup>-C<sup>10,11</sup> metallo-mediated base pairs, by incorporating Hg<sup>2+</sup> or Ag<sup>+</sup> cations, respectively. The strong metal-base interactions are highly specific in both cases, and it can interfere in the replication and transcription of DNA and induce conformational changes (i.e. from random coil conformation to duplex conformation). This highly specific conformational changes have been used as analytical probe of Ag<sup>+</sup> upon bio-recognition by C-C mismatching, by using circular dichroism spectroscopy to sense real time conformational changes<sup>12</sup> or electrochemical methods to follow the conformational-dependent activity of Exonuclease III.<sup>13</sup>

One interesting application of the strong nucleobase-metal binding feature of DNA is the production of highly fluorescent and tunable hybrid DNA-Ag<sub>n</sub> clusters. The optical properties of these hybrid systems (e.g. high fluorescence yield and absorption/emission wavelength tuning) are strongly dependent on the nucleobases sequence,<sup>14-17</sup> cluster size,<sup>18</sup> pH,<sup>19</sup> temperature<sup>20</sup> etc. These photophysical properties of silver clusters have led to the creation of a new generation of small and biocompatible fluorophores as biological

labels, that exceed the commonly used semiconductors quantum dots and organic dyes in regard to fluorescence quantum yield, photostability and biocompatibility, due to the low toxicity and very small size of the  $\text{Ag}_n$  clusters.<sup>14,21</sup> For more information about these fascinating systems the readers are recommended to refer to recent reviews on this topic.<sup>14,22 - 27</sup>

The hybrid DNA- $\text{Ag}_n$  clusters have two intense absorption bands, one in the visible which is tunable with DNA bases sequence and the other in the UV spectral region which is common to all of them, regardless of the position of the visible band.<sup>14,17,19,28</sup> The UV excitation band is located in the spectral region where the DNA bases absorb (260 – 270 nm). Very interestingly, the excitation of the common UV band leads to the same fluorescence spectrum as in the case of excitation of the tunable visible band. It was previously suggested that this peak could be due to excitation to higher lying states,<sup>28</sup> however, recent evidence indicate that the UV absorption band corresponds to excitation of the nucleobases.<sup>17</sup>

While DNA nucleobases<sup>29</sup> and protonated nucleobases<sup>30</sup> present mostly very short excited state lifetime on the sub-ps to ps time scale, highly fluorescent DNA- $\text{Ag}_n$  clusters have excited state lifetime in the ns regime,<sup>10</sup> although according to a previous report<sup>17</sup> their excitation in the UV spectral region is on the DNA moiety for which short excited state lifetimes are expected. This is clear evidence of the effect of Ag on the excited state lifetime of these systems. In addition, it has been reported that the formation of highly fluorescent DNA- $\text{Ag}_n$  clusters is favored when using C rich DNA strands as template.<sup>15,19</sup>

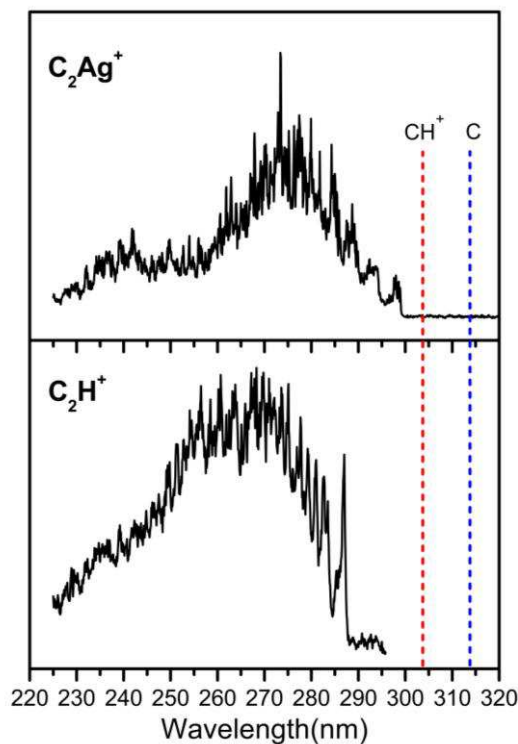
The detailed mechanism responsible for the fluorescence enhancement is still elusive and whether metal-mediated DNA base pairs will behave in a similar way as natural DNA may be a major issue to take into account when analyzing biocompatibility. In this context, gas phase characterization of optical and structural properties of model systems, together with quantum modeling, has shown to be suitable for understanding the photophysics of related hybrid aminoacids or peptides clustered with  $\text{Ag}_n^+$  or  $\text{Au}_n^+$  clusters. In previous seminal works, it has been shown that complexation of  $\text{Ag}^+$ ,<sup>31</sup>  $\text{Au}^{+32}$  and  $\text{Ag}_n^{+33-36}$  with aminoacids and small peptides significantly changes the optical properties of the system. In those cases it was manifested as a

strong absorption band around 300 – 450 nm spectral region attributed to Charge Transfer (CT) excitation. For more information on this topic, readers are referred to a very recent and complete review that deals with these systems from basics towards sensor development.<sup>37</sup>

Here we report the spectroscopic characterization in the UV region, where the nucleobase is expected to absorb, of the (Cytosine)<sub>2</sub>Ag<sup>+</sup> complex (C<sub>2</sub>Ag<sup>+</sup>) and the related protonated cluster (C<sub>2</sub>H<sup>+</sup>) for comparison of their optical properties, as a reductionist approach to gain information about the molecular mechanism that controls the appealing photophysical properties of hybrid DNA-Ag<sub>n</sub> clusters. It must be noted that although it was recently established that fluorescent DNA-Ag<sub>n</sub> clusters do contain varying amounts of cationic silver, the presence of neutral silver clusters is crucial for emission of photons following excitation.<sup>38</sup> Thus, a direct connection between the present results and fluorescent DNA-Ag<sub>n</sub> clusters is not feasible at this point and more work considering hemi-reduced silver cluster is necessary. However, it constitutes a first approach to start understanding this interesting phenomenon.

The photofragmentation spectra of C<sub>2</sub>H<sup>+</sup> and C<sub>2</sub>Ag<sup>+</sup> were recorded in the spectral range (225 – 320 nm) under similar experimental conditions for comparison are shown in Figure 1. In both cases the main fragmentation channel was the elimination of one neutral cytosine molecule.

The origins of the electronic transitions 286.9 nm for C<sub>2</sub>H<sup>+</sup> and 299.1 nm for C<sub>2</sub>Ag<sup>+</sup>, are close and also similar to the origin of the electronic transition of CH<sup>+</sup> (303.5 nm)<sup>30</sup> and C (314.2 nm).<sup>39</sup> An enlarged view of the spectra near the origin is shown in Figure 2, from which a very different vibronic structure and bandwidth can be observed for C<sub>2</sub>H<sup>+</sup> and C<sub>2</sub>Ag<sup>+</sup>. The spectrum of the C<sub>2</sub>Ag<sup>+</sup> complex shows a very low vibrational frequency progression ( $\nu = 24\text{ cm}^{-1}$ ) and another one at  $123\text{ cm}^{-1}$  which correspond to in-plane intermolecular bending modes (N-Ag<sup>+</sup>-N angle),  $\nu_2 = 33\text{ cm}^{-1}$  and  $\nu_5 = 95\text{ cm}^{-1}$  as calculated in the ground state geometry at the MP2/SV(P) theory level (see Figure S1 in the Supporting Information). It should be noted that this angle changes from 159° in the ground state optimized geometry to 165° in the A' state optimized geometry which is the main geometrical change between both equilibrium structures.



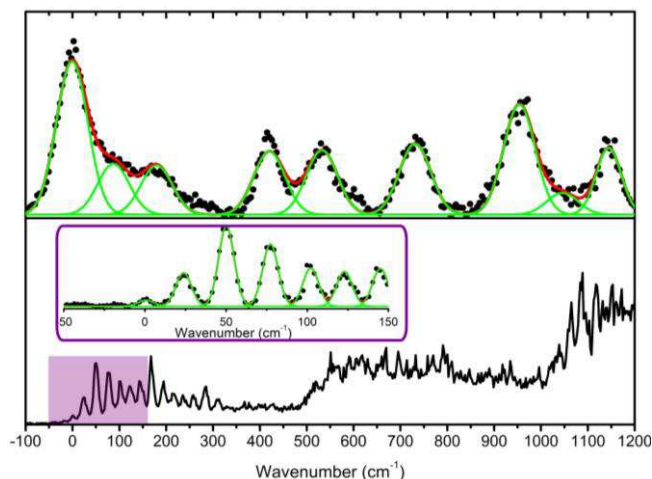
**Figure 1:** Photofragmentation spectra of the  $C_2Ag^+$  and  $C_2H^+$  complexes in the whole spectral range analyzed in this work (225 - 320) nm. Vertical dashed lines show the wavelengths of the origins of the electronic transition of the individual components of both complexes,  $C^{39}$  and  $CH^{+30}$ . The electronic transitions wavelength of Ag (328.2 nm)<sup>40</sup> and  $Ag^+$  (110.7 nm, 111.2 nm and 119.6 nm)<sup>41,42</sup> fall out of scale.

The excited state lifetimes have been estimated, as in a previous study,<sup>30</sup> from the widths of the bands fitted to Voigt profiles to account for the rotational contour and the laser Gaussian bandwidth (11  $cm^{-1}$ ) convolution. The intrinsic Gaussian profile of the laser is shown in the Figure S2 (see Supporting Information) along with the experimental profiles recorded for both complexes. The relationship between the Lorentzian full width at half maximum (FWHM) and the excited state lifetime ( $\tau$ ) is given by the uncertainty principle, which can be written as:

$$FWHM (cm^{-1}) = (2\pi c \tau)^{-1} = 5.3 \times 10^{-12} / \tau(s) \quad (\text{Eq. 1})$$

where  $c$  stands for the speed of the light. This procedure is valid if it can be assumed that the broadening is only due to the excited state lifetime and not to spectral congestion (e.g. rotational contour or low frequency active vibrational modes). The spectrum of the  $C_2Ag^+$  complex clearly shows that the spectral resolution is good enough to resolve the different vibronic transitions, even in the case of vibrational frequencies as low as  $24\text{ cm}^{-1}$ , while the rotational broadening can be neglected since the temperature of the experiment is low (40 K)<sup>30</sup>. Therefore, the broadening in the  $C_2H^+$  spectrum is due to non-radiative process leading to a short excited state lifetime. From the bandwidth analysis, the Lorentzian FWHM for the  $C_2Ag^+$  complex is less than  $1\text{ cm}^{-1}$  and can be associated to an excited state lifetime  $\tau > 5000\text{ fs}$ , while the bandwidth for the  $C_2H^+$  complex is  $FWHM = 62\text{ cm}^{-1}$  with an associated lifetime  $\tau \approx 85\text{ fs}$ , which is at least two orders of magnitude shorter than that of the former complex. This remarkable difference shows the effect of the  $Ag^+$  cation on the excited state dynamics of the complex. The long excited state lifetime estimated for the  $C_2Ag^+$  complex is compatible with high fluorescence quantum yield of DNA- $Ag_n$  clusters, with C rich oligonucleotide strands.<sup>9,10,14,16</sup> The lifetime of the  $C_2H^+$  complex is quite similar to the ones observed for the free protonated cytosine ( $133 \pm 20$ ) fs,<sup>30</sup> indicating that it is not strongly perturbed by the dimer formation.

The electronic ground state structures of the protonated cytosine dimer<sup>43,44</sup> and modified cytosine dimer,<sup>43-45</sup> and  $C_2Ag^{+46}$  in the gas phase have been previously determined by Infrared Multiphoton Dissociation (IR-MPD) spectroscopy together with quantum chemical calculations. Both complexes are planar (Cs symmetry) and have equivalent structures, in which each C molecule is found in the keto-amino form with the NH groups where the glycosidic bonds are expected, in *transoid* orientation as in the case of the *i*-motif structure of DNA. The  $H^+$  as well as the  $Ag^+$  cation acts as a bridge between the hetero-nitrogen atoms in each C (Table 1). In addition, theoretical calculations have shown that these structures are the most stable ones in the electronic ground state.<sup>43,44,46,47</sup>



**Figure 2:** Low energy part of the spectra of the  $C_2H^+$  (a) and  $C_2Ag^+$  (b) complexes. In panel (b) the inset shows an amplification with better resolution of the first 150  $cm^{-1}$  of the spectrum of the  $C_2Ag^+$  complex, in which a low vibrational frequency progression composed of narrow bands is observed. The vibrational bands were fitted to Voigt profiles.

To help the interpretation of the results and get more insight into the structure and excited state dynamics of these complexes, ground and excited state optimizations were performed at the MP2 and RI-ADC(2) (SV(P) basis set) theory levels, respectively.

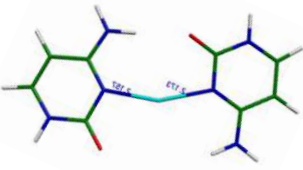
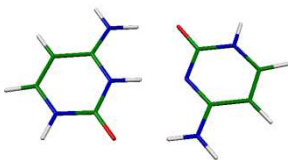
The ground state optimization at the MP2 level, lead to the same planar structures as in previous studies at other levels of theory.<sup>43,44,46,47</sup> The vertical energies ( $E_{\text{vert}}$ ) of the first excited states of  $A'$  (symmetrical versus the molecular plane) and  $A''$  (antisymmetrical versus the molecular plane) symmetry were also calculated (Table 1).

In many free neutral or protonated DNA bases, the  $S_1$  optimization leads to out-of-plane deformations and to an avoided crossing between  $S_1$  and  $S_0$  states, which is responsible for a fast non-radiative decay.<sup>48</sup> For the  $C_2H^+$  and  $C_2Ag^+$  complexes, the size and complexity of the systems imposed to limit the calculation to the accessible Franck-Condon region, so the excited states were only optimized in Cs (planar) symmetry since the ground state geometry is planar. Moreover, the  $A''$  and  $A'$  states being very close in energy, any optimization without Cs symmetry leads to calculation failure.



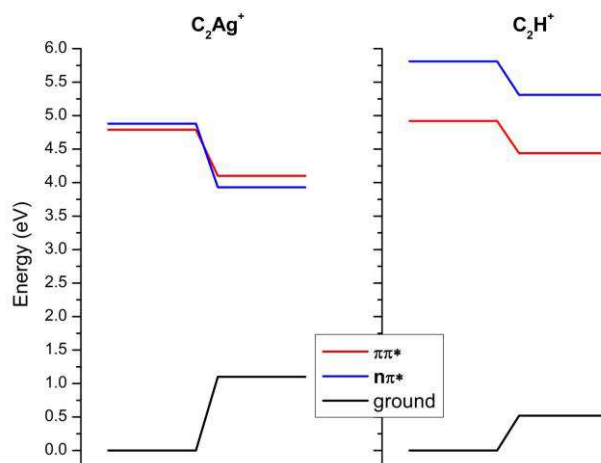
The optimization of the geometry in the  $A'$  and  $A''$  excited states allows determining the equilibrium (lowest energy) structure in each excited state and then the Cs adiabatic transition energies ( $E_{ad}$ ), which is the energy involved to reach the equilibrium geometry in the excited state from the equilibrium geometry in the ground state. Since the ground state is planar, they can be considered as the adiabatic potential accessible in the Franck–Condon window from the ground states and indeed the calculated  $E_{ad}$  values are in good agreement with the experimental transition energies ( $0-0_{exp}$ ). All the results are summarized in Table 1 and Figure 3. As usual for this type of calculations the calculated adiabatic values are within 0.2 eV from the experimental ones.<sup>30</sup>

**Table 1.** Experimental and theoretical electronic excitations energies of  $C_2Ag^+$  and  $C_2H^+$  clusters for the first  $A'$  and  $A''$  symmetry excited electronic states in planar Cs symmetry.

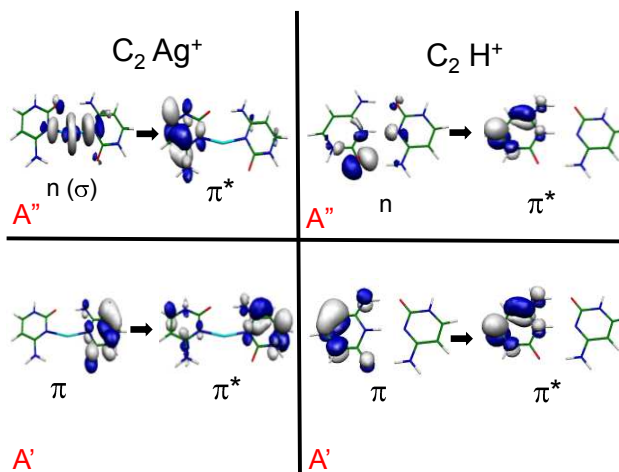
	$C_2Ag^+$ <b><math>0-0_{exp} = 4.16</math> eV</b> 			$C_2H^+$ <b><math>0-0_{exp} = 4.32</math> eV</b> 		
	$S_0^a$	$1A'(1)^a$	$1A''(1)^a$	$S_0^a$	$1A'(1)^a$	$1A''(1)^a$
Osc. Strength		0.15	0.011		0.13	0.0004
$S_0$ optimization	0.00	4.79	4.88	0.00	4.92	5.82
$A'$ optimization	1.10	<b>4.10</b>	3.93	0.52	<b>4.44</b>	5.31
$A''$ optimization	2.00	5.32	<b>3.04</b>	3.29	6.11	<b>3.63</b>

<sup>a</sup> Excitation energy values are in eV.

Highlighted in bold-black are the  $E_{ad}$  from the  $S_0$  to the first  $A''$  excited state and in bold-red those that best match with the experimental values of the 0-0 transition that correspond to the transition from the  $S_0$  to the first  $A'$  symmetry excited state in both complexes. The locally excited electronic state ( $A'$  symmetry) is a  $\pi\pi^*$  state in the  $C_2H^+$  case while for  $C_2Ag^+$ , in addition it has a small partial CT character between both cytosine molecules (Figure 4).



**Figure 3:** Scheme of the energy levels of  $C_2Ag^+$  and  $C_2H^+$ . For each complex, the left side corresponds to the vertical energies at the ground state equilibrium geometry and the right side corresponds to the energies of the ground,  $\pi\pi^*$  and  $n\pi^*$  states at the  $A'$  ( $\pi\pi^*$ ) excited state optimized structure conserving the planar geometry. In the  $C_2Ag^+$  complex the  $\pi\pi^*$  and  $n\pi^*$  states are very close in energy which might be related to the observed longer lifetime as compared with the  $C_2H^+$  complex.



**Figure 4:** Main orbitals involved in the electronic structure of the two first states of  $A'$  and  $A''$  symmetry.

In the case of the  $C_2Ag^+$  complex, at the ground state equilibrium geometry the  $\pi\pi^*$  state ( $A'$ ) is lower in energy than the first ( $A''$ )  $n_{Ag}\pi^*$  state. Upon excited state optimization of the  $A'$  states in Cs symmetry, the first  $A''$  state becomes lower in energy than the optically allowed  $A'$  state (Figure 3). We have observed a similar behavior between two tautomers of protonated uracil: the enol/enol tautomer has a short excited state lifetime while the enol/keto has a longer one.<sup>30</sup> In this latter tautomer the  $n\pi^*$  is lower in energy than the  $\pi\pi^*$  at the Cs  $\pi\pi^*$  optimized geometry as in the case of  $C_2Ag^+$ . Upon slight out-of-plane deformation, the optimization process of the “ $A$ ” state brings it back in its planar geometry, while the “ $A'$ ” state tends to be stabilized upon out-of-plane deformation. This may lead to a crossing between these states and to a barrier along the out-of-plane deformation coordinate.

On the other hand, in the case of the  $C_2H^+$  complex, the  $n_o\pi^*$  state ( $A''$ ) stays higher in energy than the  $\pi\pi^*$  ( $A'$ ) upon optimization of the  $A'$  states in Cs symmetry. In absence of Cs symmetry both states are mixed and optimization of the  $S_1$  state leads to a crossing with the ground state. Maybe, there is a correlation between the energy gap between the  $\pi\pi^*$  and the  $n\pi^*$ , the lifetime being longer when the  $n\pi^*$  is slightly lower in energy at the  $\pi\pi^*$  optimized geometry than when the  $n\pi^*$  is higher in energy. This should be tested theoretically but on simpler systems which less spectral congestion.

One may tentatively rationalize the variation of the excited state lifetime in the following way. The optical excitation prepares the  $\pi\pi^*$  state in its Cs optimized geometry, i.e. planar or nearly planar. If there is no  $n\pi^*$  state lower in energy, the  $S_1$   $\pi\pi^*$  state will easily undergo out-of-plane deformations, which leads to internal conversion. At the opposite if the  $n\pi^*$  is lower in energy than the  $\pi\pi^*$  state, out-of-plane deformations will induce a crossing between these states, resulting in a barrier for the  $\pi\pi^*$  state along the out-of-plane coordinate that will prevent the second crossing with the ground state at low excitation energies. Thus this state should have a longer lifetime.

This idea has to be tested by more elaborate calculations on this system or even on the simplest protonated DNA bases for which there is now some experimental information.<sup>30</sup>

In summary, the complexation of two cytosine molecules with  $\text{Ag}^+$  does not change drastically the character and energy of the electronic transition as compared to  $\text{C}_2\text{H}^+$  or  $\text{CH}^+$ , contrary to the silver/gold aminoacids case, for which a new absorption band appears in the near UV/visible spectral region (300 – 450 nm), depending on the specific system, attributed to CT excitations.<sup>31 - 37</sup> The present results are in agreement with previous results<sup>17</sup> that suggested that, in bulk systems, the UV excitation of highly fluorescent DNA- $\text{Ag}_n$  clusters is due to absorption of the DNA bases. On the other hand, the  $\text{Ag}^+$  complexation significantly changes the excited state lifetime of the complex as compared with  $\text{H}^+$  complexation. This change is explained as a consequence of the removal of a conical intersection between the excited and the ground states of  $\text{C}_2\text{Ag}^+$  complex, which takes place in the  $\text{C}_2\text{H}^+$  complex and leads to a fast non-radiative decay of the excited state of the latter. Therefore, a higher fluorescence quantum yield is expected upon  $\text{Ag}^+$  complexation and this could be the reason for the high fluorescence of DNA- $\text{Ag}_n$  clusters, although in the latter case the fluorescence is observed upon chemical reduction of  $\text{Ag}^+$ . However, it has been shown that cationic systems can give relevant and detailed information to understand the optical properties of hybrid biomolecules-noble metal clusters.<sup>31 -</sup>

37

The ultimate goal of noble-metal bioconjugation is to develop biosensing at the molecular level and gas phase studies can help to build model systems in a bottom-up strategy. In this regard, the next step will be to generate larger  $\text{Ag}_n^+$  clusters conjugated with DNA bases, in which some reduction processes has taken place.

## Methodology

### a. Experiment

The electronic spectra of the  $\text{C}_2\text{Ag}^+$  and  $\text{C}_2\text{H}^+$  complexes were obtained via parent ion photo-fragment spectroscopy in a cryogenically-cooled quadrupole ion trap (Paul Trap from Jordan TOF Products, Inc.).<sup>49</sup> The setup is similar to the one developed in several groups based on the original design by Wang and Wang.<sup>50-</sup>

<sup>52</sup>The complexes are produced in an electrospray ionization source built at Aarhus University,<sup>53</sup> by introducing

a solution of cytosine (500  $\mu\text{M}$ ) and silver nitrate (250  $\mu\text{M}$ ) in a methanol (50%)/water (50%) solvent . At the exit of the capillary, ions are trapped in an octopole trap for 90 ms. They are extracted by applying a negative pulse of c.a. 50 V and are further accelerated to 190 V by a second pulsed voltage just after the exit electrode. This time sequence of pulsed voltages produces ion packets with duration between 500 ns and 1  $\mu\text{s}$ . The ions are driven by a couple of electrostatic lenses toward the Paul trap biased at 190 V so that the ions enter the trap gently avoiding fragmentation induced by collisions. A mass gate placed at the entrance of the trap allows selecting the parent ion. The Paul trap is mounted on the cold head of a cryostat (Coolpak Oerlikon) connected to a water-cooled He compressor. Helium as buffer gas is injected in the trap using a pulsed valve (General Valve) triggered 1 ms before the ions enter the trap as previously reported by Kamrath et al.<sup>51</sup> The ions are trapped and thermalized at a temperature between 20 and 50 K through collisions with the cold buffer gas. The ions are kept in the trap for several tens of ms before the photodissociation laser is triggered. This delay is necessary to ensure thermalization of ions and efficient pumping of the He buffer gas from the trap to avoid collision induced dissociation of the ions during the extraction towards the 1.5 m long time-of-flight mass spectrometer. After laser excitation, the ions are stored in the trap for a delay that can be varied between 20 and 90 ms before extraction to the TOF mass spectrometer. The complete mass spectrum is recorded on a micro channel plate (MCP) detector with a digitizing storage oscilloscope interfaced to a PC. The photofragment yield spectrum of each detected ion is normalized to the parent ion signal and the laser power. The photo-dissociation laser is an OPO laser from EKSPLA, which has a 10 Hz repetition rate, 10 ns pulse width, a resolution of 10  $\text{cm}^{-1}$  and a scanning step of 0.02 nm. The laser is shaped to a 1mm<sup>2</sup> spot to fit the entrance hole of the trap and the laser power is around 20 mW in the UV spectral region.

## **b. Calculations**

*Ab initio* calculations have been performed with the TURBOMOLE program package,<sup>54</sup> making use of the resolution-of-the-identity (RI) approximation for the evaluation of the electron-repulsion integrals.<sup>55</sup> The

equilibrium geometry of the clusters and the vibrational frequencies in their ground electronic state ( $S_0$ ) were determined at the MP2/ SV(P) level. Excitation energy and equilibrium geometry of the lowest excited singlet state ( $S_1$ ) were determined at the RI-ADC(2)/SV(P) level.

Structure optimizations were done by the quasi-Newton Raphson methods using the exact gradient vector and an approximation to the Hessian matrix as implemented in the TURBOMOLE program package.

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**Supporting Information Available:** Description and assignment of the low frequency vibrational active modes of the  $C_2Ag^+$  cluster, and the intrinsic Gaussian profile of the laser together with the experimental profiles recorded for both complexes. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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